IN THE CLAIMS:

The following listing of claims will replace all prior versions and listings of the claims in the present application:

1. (Original) An isolated polynucleotide comprising a regulatory region containing a nucleotide sequence less than about 1000 nucleotides long selected from the group consisting of:

(a) SEQ ID NO: 16, a fragment of SEQ ID NO: 16, the complement of SEQ ID NO: 16, or a fragment of the complement of SEQ ID NO: 16;

(b) a polynucleotide that hybridizes to the polynucleotide of (a) under conditions of high stringency; and

(c) a polynucleotide with at least 80% sequence homology to the polynucleotide of (a).

2. (Original) An isolated polynucleotide comprising a regulatory region containing a nucleotide sequence less than about 1000 nucleotides long selected from the group consisting of:

(a) SEQ ID NO: 17, a fragment of SEQ ID NO: 17, the complement of SEQ ID NO: 17, or a fragment of the complement of SEQ ID NO: 17;

(b) a polynucleotide that hybridizes to the polynucleotide of (a) under conditions of high stringency; and

(c) a polynucleotide with at least 80% sequence homology to the polynucleotide of (a).

3. (Original) An isolated polynucleotide comprising a regulatory region containing a nucleotide sequence less than about 1000 nucleotides long selected from the group consisting of:

(a) SEQ ID NO: 18, or a fragment of the complement of SEQ ID: 18, the complement of SEQ ID NO: 18, or a fragment of the complement of SEQ ID NO: 18;

(b) a polynucleotide that hybridizes to the polynucleotide of (a) under conditions of high stringency; and

(c) a polynucleotide with at least 80% sequence homology to the polynucleotide of (a).

4. (Original) An isolated polynucleotide comprising a regulatory region containing a

nucleotide sequence less than about 1000 nucleotides long selected from the group consisting of:

(a) SEQ ID NO: 19, a fragment of SEQ ID NO: 19, the complement of SEQ ID NO: 19, or a fragment of the complement of SEQ ID NO: 19;

(b) a polynucleotide that hybridizes to the polynucleotide of (a) under conditions of high stringency; and

(c) a polynucleotide with at least 80% sequence homology to the polynucleotide of (a).

5. (Original) An isolated polynucleotide comprising a regulatory region containing a nucleotide sequence less than about 1000 nucleotides long selected from the group consisting of:

(a) SEQ ID NO: 20, a fragment of SEQ ID NO: 20, the complement of SEQ ID NO: 20, or a fragment of the complement of SEQ ID NO: 20;

(b) a polynucleotide that hybridizes to the polynucleotide of (a) under conditions of high stringency; and

(c) a polynucleotide with at least 80% sequence homology to the polynucleotide of (a).

6. (Original) An isolated polynucleotide comprising a regulatory region containing a nucleotide sequence less than about 1000 nucleotides long selected from the group consisting of:

(a) SEQ ID NO: 21, a fragment of SEQ ID NO: 21, the complement of SEQ ID NO: 21, or a fragment of the complement of SEQ ID NO: 21;

(b) a polynucleotide that hybridizes to the polynucleotide of (a) under conditions of high stringency; and

(c) a polynucleotide with at least 80% sequence homology to the polynucleotide of (a).

7. (Previously presented) A recombinant vector comprising the isolated polynucleotide of any of claims 1-6 or 23-24, operably linked to a heterologous coding region.

8. (Previously presented) An expression cassette comprising operably linked in 5' to 3' order the isolated polynucleotide of any of claims 1-6 or 23-24, a heterologous coding region, and a termination sequence.

9. (Original) A host cell comprising the vector of claim 7.

10. (Original) The host cell of claim 9 wherein the host cell is a yeast cell.

11. (Original) The yeast cell of claim 10 wherein the yeast cell is a methylotrophic yeast cell.

12. (Original) The methylotrophic yeast cell of claim 11 wherein the yeast cell is selected from the group of genera consisting of *Hansenula*, *Candida*, *Torulopsis*, and *Pichia*.

13. (Original) The yeast cell of claim 12 wherein the yeast cell is from *Pichia pastoris*.

14. (Original) A host cell comprising the expression cassette of claim 8.

15. (Original) The host cell of claim 14 wherein the host cell is a yeast cell.

16. (Original) The host cell of claim 15 wherein the yeast cell is a methylotrophic yeast cell.

17. (Original) The host cell of claim 16 wherein the methylotrophic yeast cell is selected from the group of genera consisting of *Hansenula*, *Candida*, *Torulopsis* and *Pichia*.

18. (Original) The host cell of claim 17 wherein the yeast cell is from *Pichia pastoris*.

19. (Original) The host cell of claim 9 wherein the host cell expresses a protein encoded by the vector.

20. (Original) The host cell of claim 14 wherein the host cell expresses a protein encoded by the expression cassette.

21. (Original) A method for the production of a protein comprising growing the host cells of claim 19 under conditions where the host cells express the protein encoded by the vector and isolating the expressed protein.

22. (Original) A method for the production of a protein comprising growing the host cells of claim 20 under conditions where the host cells express the protein encoded by the vector and isolating the expressed protein.

23. (Previously presented) An isolated polynucleotide comprising the nucleotide sequence as set forth in SEQ ID NO: 30.

24. (Previously presented) An isolated polynucleotide comprising the nucleotide sequence as set forth in SEQ ID NO: 31.

25. (New) An isolated nucleic acid comprising SEQ ID NO: 16 and at least one regulatory element selected from the group consisting of SEQ ID NO: 17, SEQ ID NO: 18, SEQ ID NO: 19, SEQ ID NO: 20, and SEQ ID NO: 21, wherein said at least one regulatory element is placed in said nucleic acid in relation to SEQ ID NO: 16 in a non-naturally occurring manner.

26. (New) The isolated nucleic acid of claim 1, comprising multiple copies of SEQ ID NO: 16.

27. (New) The isolated nucleic acid of claim 1, comprising multiple copies of SEQ ID NO: 17.

28. (New) The isolated nucleic acid of claim 1, comprising multiple copies of SEQ ID NO: 18.

29. (New) The isolated nucleic acid of claim 1, comprising multiple copies of SEQ ID NO: 19.

30. (New) The isolated nucleic acid of claim 1, comprising multiple copies of SEQ ID NO: 20.